SEQUENCE LISTING <110> Cobb, Melanie Hutchison, Michele Chen, Zhu Berman, Kevin <120 TAO PROTEIN KINASE POLYPEPTIDES AND METHODS OF USE THEREFOR <130> **\6**0098.421C1 <150> 09****060,410 <151> 19**9**8-04**-**14 <160> 28 <170> FastSAQ for Windows Version 4.0 <210> 1 <211> 3312 <212> DNA <213> Rattus norvegicus <220> <221> CDS <222> (121)...(312**3**) <400> 1 tctgcagtat ggtagattat tatttatgca tttatgccag tgtggcttca ttcatacaga tgaaccaagc tttgggatag\cagtataaaa ttagaatcag acagctgact gctcagcagg atg cca tca act aac aga gca ggc agt cta aag gac cct gaa atc gca 168 Met Pro Ser Thr Asn Arg\Ala Gly Ser Leu Lys Asp Pro Glu Ile Ala 1 216 gag ctc ttc ttc aaa gaa ga $oldsymbol{ol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{ol}oldsymbol{ol}oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{oldsymbol{ol}}}}}}}}}}}}}}$ Glu Leu Phe Phe Lys Glu Asp\Pro Glu Lys Leu Phe Thr Asp Leu Arg 20 25 gaa atc ggc cat ggg agc ttt gga gca gtt tat ttt gca cga gat gtg Glu Ile Gly His Gly Ser Phe Gly Ala Val Tyr Phe Ala Arg Asp Val 264 35 312 cgt act aat gaa gtg gtg gcc atc aag\aaa atg tct tat agt gga aag Arg Thr Asn Glu Val Val Ala Ile Lys $oldsymbol{\chi}_{i}$ ys Met Ser Tyr Ser Gly Lys 50 5.5 360 cag tot act gag aaa tgg cag gat att att\aag gaa gtc aag ttt cta Gln Ser Thr Glu Lys Trp Gln Asp Ile Ile Lys Glu Val Lys Phe Leu 65 408 caa aga ata aaa cat ccc aac agt ata gaa tac\aaa ggc tgc tat tta Gln Arq Ile Lys His Pro Asn Ser Ile Glu Tyr Lys Gly Cys Tyr Leu 85 90 456 cgt gaa cac aca gca tgg ctt gta atg gaa tat tgt\tta gga tct gct

TO410

4

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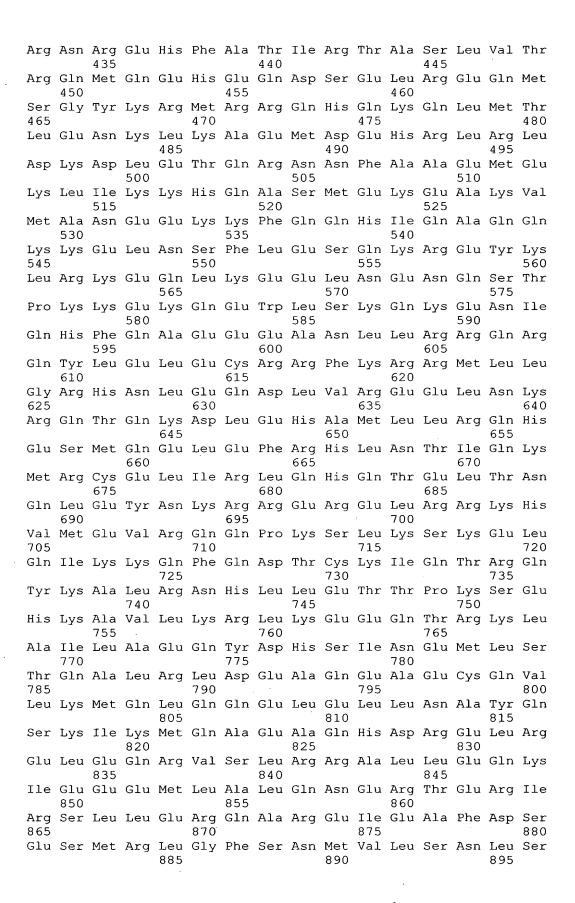
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Pro	Glu	Ala	Phe 900	Ser	His	Ser	Tyr	Pro 905	Gly	Ala	Ser	Ser	Trp 910	Ser	His	
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?

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Ile Ile His Arg Leu Pro Gly Ser Asp Asn Leu Tyr Asp Asp Pro Tyr Gln Pro Glu Met Thr Pro Gly Pro Leu Gln Pro Pro Ala Ala Pro Pro Thr Ser Thr Ser Ser Ser Ala Arg Arg Ala Tyr Cys Arg Asn Arg Asp His Phe Ala Thr Ile Arg Thr Ala Ser Leu Val Ser Arg Gln Ile Gln Glu His Glu Gln Asp Ser Ala Leu Arg Glu Gln Leu Ser Gly Tyr Lys Arg Met Arg Arg Gln His Gln Lys Gln Leu Leu Ala Leu Glu Ser Arg Leu Arg Gly Glu Arg Glu Glu His Ser Gly Arg Leu Gln Arg Glu Leu Glu Ala Gln Arg Ala Gly Phe Gly Thr Glu Ala Glu Lys Leu Ala Arg Arg His Gln Ala Ile Gly Glu Lys Glu Ala Arg Ala Ala Gln Ala Glu Glu Arg Lys Phe Gln Gln His Ile Leu Gly Gln Gln Lys Lys Glu Leu Ala Ala Leu Leu Glu Ala Gln Lys Arg Thr Tyr Lys Leu Arg Lys Glu Gln Leu Lys Glu Glu Leu Gln Glu Asn Pro Ser Thr Pro Lys Arg Glu Lys Ala Glu Trp Leu Leu Arg Gln Lys Glu Gln Leu Gln Gln Cys Gln Ala Glu Glu Ala Gly Leu Leu Arg Arg Gln Arg Gln Tyr Phe Glu Leu Gln Cys Arg Gln Tyr Lys Arg Lys Met Leu Leu Ala Arg His Ser Leu Asp Gln Asp Leu Leu Arg Glu Asp Leu Asn Lys Lys Gln Thr Gln Lys Asp Leu Glu Cys Ala Leu Leu Leu Arg Gln His Glu Ala -685 Thr Arg Glu Leu Glu Leu Arg Gln Leu Gln Ala Val Gln Arg Thr Arg Ala Glu Leu Thr Arg Leu Gln His Gln Thr Glu Leu Gly Asn Gln Leu Glu Tyr Asn Lys Arg Arg Glu Gln Glu Leu Arg Gln Lys His Ala Ala Gln Val Arg Gln Gln Pro Lys Ser Leu Lys Val Arg Ala Gly Gln Leu Pro Met Gly Leu Pro Ala Thr Gly Ala Leu Gly Pro Leu Ser Thr Gly Thr Leu Ser Glu Glu Gln Pro Cys Ser Ser Gly Gln Glu Ala Ile Leu Gly Gln Arg Met Leu Gly Glu Glu Glu Ala Val Pro Glu Arg Met Ile Leu Gly Lys Glu Gly Thr Thr Leu Glu Pro Glu Glu Gln Arg Ile Leu Gly Glu Glu Met Gly Thr Phe Ser Ser Pro Gln Lys His Arg Ser Leu Val Asn Glu Glu Asp Trp Asp Ile Ser Lys Glu Met Lys Glu Ser Arg Val Pro Ser Leu Ala Ser Gln Glu Arg Asn Ile Ile Gly Gln Glu Glu Ala Gly Ala Trp Asn Leu Trp Glu Lys Glu His Gly Asn Leu



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                           1145
                                            1150
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                                         1165
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                                      1180
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                                  1195
Pro Arg Ser Gln Arg Arg Leu Gly Leu Ser Ala Ser Arg Gln Leu Pro
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			20				_	25					30			
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	50					55	_	_			60		-	Lys		
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225					230					235				Trp	240	
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ьeu	$\sigma \perp u$	rne	νdΤ	GTII	rne	тт6	HSD	ьуз	Cys	neu	HT A	шγз	ETO	Ala	GIU	

270

265

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Gly Thr Asn Val Ser Val Ala Ile Lys Gln Met Asn Leu Glu Lys Gln
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Lys His Pro Asn Ile Val Asn Phe Ile Asp Ser Tyr Val Leu Lys Gly
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Asp Leu Trp Val Ile Met Glu Tyr Met Glu Gly Gly Ser Leu Thr Val
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Cys Arg Glu Thr Leu Ser Gly Leu Glu Phe Leu His Ser Lys Gly Val
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Leu His Arg Asp Ile Lys Ser Asp Asn Ile Leu Leu Ser Met Glu Gly
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Asn Leu Lys Arg Thr Thr Met Val Gly Thr Pro Tyr Trp Met Ala Pro
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Leu Gly Ile Met Ile Ile Glu Met Ile Glu Gly Glu Pro Pro Tyr Leu
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Asn Glu Thr Pro Leu Arg Ala Leu Tyr Leu Ile Ala Thr Asn Gly Thr
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Pro Lys Leu Lys Glu Pro Glu Asn Leu Ser Ser Leu Lys Lys Phe
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35

392

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25

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						gga Gly										776
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		_	_			-								aaa Lys		2840
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Glu														cgt Arg		3080
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Ile Gln His Leu Asn Glu Leu His Ala Met Lys Lys Arg His Leu 🛭 🛭 🖽 Thr Gln His Glu Ala Glu Ser Ala Ser Gln Asn Glu Tyr Thr Gln Arg Gln Gln Asp Glu Leu Arg Lys Lys His Ala Met Gln Sex Arg Gln Gln 780/ Pro Arg Asp Leu Lys Ile Gln Glu Ala Gln Ile Arg Lys Gln Tyr Arg Gln Val Val Lys Thr Gln Thr Arg Gln Phe Lys Leu Tyr Leu Thr Gln Met Val Gln Val Val Pro Lys Asp Glu 🗹 Lys Glu Leu Thr Ser Arg Leu Lys Gln Asp Gln Met Gln Lys 🌿 Ala Leu Leu Ala Ser Gln Tyr Glu Ser Gln Ile Lys Lys Met Val Gln Asp Lys Thr Val Lys Leu Glu Ser Trp Gln Glu Asp Glu Gln Arg Val Leu Ser Glu Lys Leu Glu Lys Glu Leu Glu Glu Leu I 🖟 Ala Tyr Gln Lys Lys Thr Arg Ala Thr Leu Glu Glu Gln Ile Ly& Lys Glu Arg Thr Ala Leu Glu Glu Arg Ile Gly Thr Arg Arg Ala Met Leu Glu Gln Lys Ile Ile Glu Glu Arg Glu Gln Met Gly Glu Met Arg Arg Leu Lys Lys Glu Gln Ile Arg Asp Arg His Ser Gln Glu Arg His Arg Leu Glu Asn His Phe Val Arg Thr Gly Ser Thr Ser/Arg Ser Ser Gly Gly Ile Ala Pro Gly Val Gly Asn Ser Ser Ser The Gln Met Ala Met